

Fig. 1

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Y VLOL YISLNSDMYERY VLOL YISLNSDMYERY VLOI IITVNSDLVEN N YLAM IICOSAAAFEN N YISM IILVOTAAGFEN YLAM VLAL VLRTTATGRI SV N IATEGSLI LOSV S IKIENTLI LOSV S IKIENTLI LOGV S IKIENTLI LOGV S IKTENKLI POW S IK	SLKITWLGGKQVVNVIIRVNDYLSERPKIRVTTIPITQNYLGASGRLLKLGDRVXINT ALKITWLKKRQVVNVLIRINNYLSDRPKIVVETIPITQNYLGASGRLLKLGKKIYIIT ASHSPWFSDRRWVNSIIVVDKGLNSIPKLKVWTISMRQNYWGSEGRLLLLGGRKIYIIT AKSSYXSWFGNRMIQSGILACPLRQDLTNECLVLPFSNDQVLMGAEGRLYMYGDSVYYYQ ARSSYVIRYHSNRLLQSAVLICPLSDMHTARCNLVMFNNSQVMMGAEGRLYVIDNNYYYQ ALRSYFPSYFSNRRVQSAFLVCAWNQILVTRCELVVPSNNQTLMGAEGRLYVLINNRLLYYQ AKSSYKPGRFGGKRIQQAILSIKVSTSLGEDPVLTVPPNTVTLMGAEGRLYVLINNRLLYYQ AKSSYKPGRFGGKRIQQAILSIKVSTSLGEDPVLTVPPNTVTLMGAEGRLLTVGTSHFLYQ AKSSYKPGRFGGKRIQQAILSIKVSTSLGEDPVLTVPPNTVTLMGAEGRLLTVGTSHFLYQ ANSSYRPGAEGRFGGKRIQQAILSIKVSTSLGEDPVLTVPPNTVTLMGAEGRLLTVGTSHFLYQ ARSSYKPGRFGGKRIQQAILSIKVSTSLGEDPVLTVPPNTVTLMGAEGRLLTVGTSHFLYQ ARSSYKPGRFGGKRIQQAILSIKVSTSLGEDPVLTVPPNTVTLMGAEGRLLTVGTSHFLYQ ARSSYKPGRFGGKRIQAARST	WATUT YANTSOV TIM VATUT YANTSOV TIM VEGSEATFIGSY NIATO I TMY ONALNPNYRFAGAF RNESN T TFY CINRNFYFIGAL NSSTT V TLY LRGVFGTM DSEQA L ASA B5S2	
L TOG B L TOG B I V LNG C I V LNG C I V LGD C I V LGD C I V LGD C I V LSG C I V	SERPKIRVTTIPI SDRPKIVVETIPI NSIPKLKVWTISMI DLTNECLVLPFSNI MHTARCNLVMFNN SLGEDPVLTVPPN BASS4	AAAN TGSI PSSTTS.	IPKLCKAES IPKICKITS IPKSCS VTLS LIP LITI ILKNDGVREARSG
IS V L LSIGEAIYA SS JU V L LSIGDAIXA SS JU V I LVINDLIYA TS PE T I FSLIKTHWC TH PN T I FSLIKTHWC TH GS T I FDMSATHYC TH GS T I FDMSATHYC TH BISI BISI EDLVLDVLDLKGSTKSHRYR] EDLVLDLNHDGSISTTRFK) PEQRIIIMYNDTIVERII. AELRLAFYYNDTIVERII. PTQKLTLLFYNDTVTERII. TLMAHGRLGFDGQYHEKDL. FRANALASS	GGKQVVNVIIRVNDYLSERPKIRVTTI KKRQVVNVLIRINNYLSDRPKIVVETI SDRRMVNSIIVVDKGLNSIPKLKVWTI GNRMIQSGILACPLRQDLTNECLVLPF SNRLLQSAVLICPLSDMHTARCNLVWF SNRRVQSAFLVCAWNQILVTNCELVVP GGKRIQQAILSIKVSTSLGEDPVLTVP	IS TAYP SPI IS TAYP SPI IT TAYP NPI IT A AWL TNDI VS T PWP TPI VT T PYP IF	INOKSINTLOPMLEKTSIPKLCKAB INOTSINTLOPMLEKTSIPKICKIT INHKSLDTFQPMLEKTEIPKSCS ISSLLGQFQIVPFIRQVTLS MGSSLLGEFQILPFLRELIP LASNIVGEFQILPVLTRITT ISNTLFGEFRIVPLLVEILKNDGVR
SGS TIS V L I AMP TVD V T I PTA TPE T I E PSA SPN T I E PAP TGS T I I PAP TGS T I I BAP TGS T I I BAP TGS T I I BASGIEDLVLDVLDI SSEGIEDLVEDIVLDI SSEGIEDLVEDIVLDI SSEGIEDLVEDIVLDI NSAVPTLAAHGRLG NSAVPTLAAHGRLG NSAVPTLAAHGRLG	TWL TWL SXWE SXWE	E NWYNT KE D NWYNR RE E PWGHS DG D SATINR OF NATSF N SGENV TA	
LLLGPSLLL LNPGPSLLL LNNPSFI LLNNPSFI LLNNPSFI PLNMPSFI PLNMPSFI POTUN TT PTVD RT PTVD RT PTVD RT PTVD RT PTVD RT TT T	QOVSQDTCNEELKI ANVNQSVCNDELKI PGKTQRDCNQASHS SQNQATQVQNAKSSYVJAGKSSEQAAAARSSYVJ SGPQQDLDQRALRSYFF PDEQDYQIRMAKSSYKI		ITHEGRG EH ITHEGRG FH ITHYNKG FH FRDIGSVMV IY FKNTGTQKI LI FQDIGDASV VY FKVVKTNKT LS BKVVKTNKT LS
	COOCONTRANO COOCONTRANO COOCONTRANO FIRMANIA FIRMANIA FREVINE SGE VIYKRYNDIA	H. PLTINW N. PMTIKW DYSDIRIKW QPSAISAON IPPIIEAOW GOSSVNMSW NKTATLHSP	A HHHHS S HHHHS S G H HHHHS S G HHHHHS S G HHHHH S G HHHHH S G HHHHH S G H HHHH S G S G
TIAVHHAEG APLEPHSFWRCPV(RIT. HDVG KPLNPDDFWRCPV(RIT. HDVG KPLNPDDFWRCTS(AALINDNRY NGINKFIVENSYN(IPLVNDLRF NGINKFILEDYAT 1PLVNDLRF GGIGKELIVDNAS: 24) GAPIHDPDF GGIGKELIVDNAS: B5S0 DLNPVVSHTYDIND DLNPVSHTYDIND DLNPRISHTKNIND FFRTIKTLYLSDGY IFRTIKTLYLSDGY IFRTIKTLYLSDGY VAA B1S4	LITPLOGDIKCRI LITPLOGDIKCVI COLEHPINENAICNI COLEHPINENAICNIKY COLIKGTSIWNNOANKY COLIKGTPSYNNOSRY COLIKGTPSYNSAREF COLIKPNSPSDIVQEGKY RASS.		YSNTTNIINMIRIKDVQLEV. YSNTSEIINMIRIKNVQLEA. YSTATERVNELAIRNKTLSA. IANNTQIISSQOEG. TASASALLNTTGFNNTNHKA. VSALNNLKVLAPYGNQGLFA. 5) VFDSTSRSRITRVSSSSTKA.
Sendai PIV1 PIV3 SV5 MUMPS MUMPS NDV (124) Sendai PIV1 PIV3 SV5 PIV3 SV5 MUMPS	Sendai PIV1 PIV3 SV5 PIV2 Mumps	Sendai PIV1 PIV3 SV5 PIV2 Mumps NDV (416)	Sendai PIV1 PIV3 SV5 SV5 PIV2 MUMPS

Fig. 3

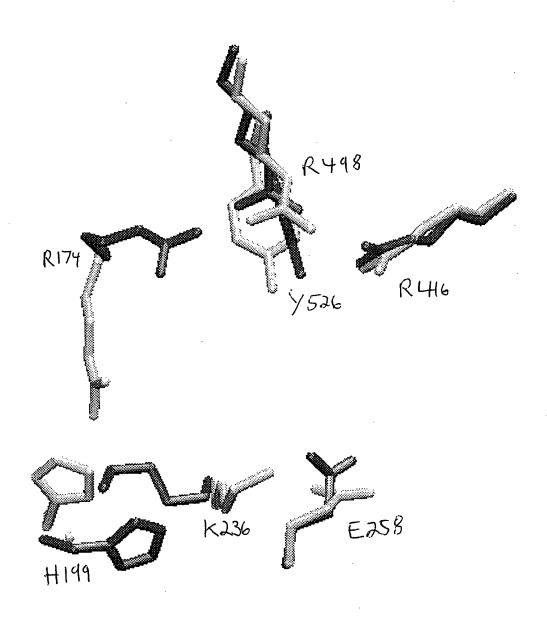


Fig. 4

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	Native 1	Native 2	Native 3	Native 4	NANA	DANA
Cell (Å)	73.3	72.3	71.7	72.0	71.6	137.5
	78.0	77.9	77.9	83.8	77.6	137.5
	202.6	199.2	198.2	201.5	197.2	116.6
Temp(K), pH	293, 7	100, 4.6	100, 4.6	293, 6	100K, 4.6	100K,6.5
Resolution	3.0	2.0	2.5	3.0	2.5	2.8
# obs	172104	623166	420703	277932	210555	498619
# unique	20022	68217	38168	22207	29671	38673
Complete ness(%)	83	86	97	88	76	94
R _{merg} (%)	9.3	4.9	3.1	9.3	4.2	5.2
R-factor			0.222		.223	0.209
R _{free}			0.277	:	.291	0.235
# protein atoms			6914		6914	6896
# CHO, Ca,			89		116	111

Fig. 5



ligands

# waters	211	207	239
 A, B Å²	25,36	32,44	44,44
 ligand Å²		38,57	48,41

Table 1 Crystallographic data and refinement statistics.

Datasets Native2, Native3, NANA and DANA from frozen crystals were collected on beamlines 11 and BW7A at DESY, Hamburg. All other datasets were collected on in-house rotating anode and image plate or multiwire detector systems.

Remerge = $\sum_{hkl} \sum_{i} |\sum_{hkl} i^{i}_{hkl} - \langle i^{i}_{hkl} \rangle |/\sum_{hkl} \sum_{i} \langle i^{i}_{hkl} \rangle$ where the sum i is over all separate reasurements of the unique reflections hkl.

$$R$$
-factor = Σ_{hkl} $||F_{obs}| - |F_{calc}|| / Σ_{hkl} $|F_{obs}|$$

R_{free}, as R-factor but summed over a 10% test set of reflections.